

## 2-Carboxyanilinium chloride monohydrate

Syed Azeem Rashid Ali Zaidi,<sup>a</sup> M. Nawaz Tahir,<sup>b\*</sup> Javed Iqbal<sup>a</sup> and Muhammad Ashraf Chaudhary<sup>c</sup>

<sup>a</sup>Institute of Chemistry, University of the Punjab, Lahore 54590, Pakistan,

<sup>b</sup>Department of Physics, University of Sargodha, Sargodha, Pakistan, and

<sup>c</sup>Department of Chemistry, F.C. College & University, Lahore, Pakistan

Correspondence e-mail: dmntahir\_uos@yahoo.com

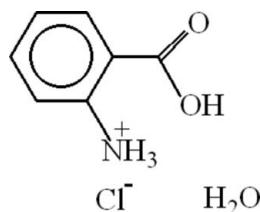
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Key indicators: single-crystal X-ray study;  $T = 296\text{ K}$ ; mean  $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$ ;  $R$  factor = 0.036;  $wR$  factor = 0.102; data-to-parameter ratio = 16.1.

In the molecule of the title compound,  $\text{C}_7\text{H}_8\text{NO}_2^+\cdot\text{Cl}^-\cdot\text{H}_2\text{O}$ , an intramolecular  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bond results in the formation of a non-planar six-membered ring adopting a flattened boat conformation. In the crystal structure, intermolecular  $\text{O}-\text{H}\cdots\text{O}$  and  $\text{N}-\text{H}\cdots\text{Cl}$  hydrogen bonds link the molecules. There is a  $\text{C}=\text{O}\cdots\pi$  contact between the carbonyl unit and the centroid of the benzene ring. There is a  $\text{C}=\text{O}\cdots\pi$  contact [ $\text{C}\cdots\text{Cg} = 3.5802(18)$ ,  $\text{C}-\text{O}\cdots\text{Cg} = 89(1)^\circ$ ] between the carbonyl unit and the centroid of the benzene ring.

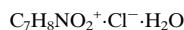
### Related literature

For applications of anthranilic acid derivatives, see: Congiu *et al.* (2005); Nittoli *et al.* (2005). For a related structure, see: Bahadur *et al.* (2007); For bond-length data, see: Allen *et al.* (1987). For ring puckering parameters, see: Cremer & Pople (1975).



### Experimental

#### Crystal data



$M_r = 191.61$

Monoclinic,  $C2/c$

$a = 23.094(4)\text{ \AA}$

$b = 4.7833(8)\text{ \AA}$

$c = 16.381(3)\text{ \AA}$

$\beta = 91.605(9)^\circ$

$V = 1808.8(5)\text{ \AA}^3$

$Z = 8$

Mo  $K\alpha$  radiation

$\mu = 0.39\text{ mm}^{-1}$

$T = 296(2)\text{ K}$

$0.28 \times 0.10 \times 0.06\text{ mm}$

#### Data collection

Bruker Kappa APEXII CCD

diffractometer

Absorption correction: multi-scan

(*SADABS*; Bruker, 2005)

$T_{\min} = 0.975$ ,  $T_{\max} = 0.985$

9747 measured reflections

2238 independent reflections

1749 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.025$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.036$

$wR(F^2) = 0.101$

$S = 1.05$

2238 reflections

139 parameters

H atoms treated by a mixture of independent and constrained refinement

$\Delta\rho_{\max} = 0.30\text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.22\text{ e \AA}^{-3}$

**Table 1**  
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O1—H1 $\cdots$ O3 <sup>i</sup>	0.82 (2)	1.73 (2)	2.539 (2)	171 (2)
N1—H1A $\cdots$ O2	0.91 (2)	1.91 (2)	2.6820 (19)	142.1 (18)
N1—H1B $\cdots$ Cl1 <sup>ii</sup>	0.88 (2)	2.28 (2)	3.1464 (16)	166.6 (18)
N1—H1C $\cdots$ Cl1 <sup>iii</sup>	0.958 (19)	2.181 (19)	3.1231 (16)	167.5 (18)
O3—H3A $\cdots$ Cl1	0.76 (3)	2.42 (3)	3.1452 (18)	160 (3)
O3—H3B $\cdots$ O2 <sup>iv</sup>	0.78 (3)	2.04 (3)	2.777 (3)	159 (3)

Symmetry codes: (i)  $-x + \frac{1}{2}, y + \frac{1}{2}, -z + \frac{1}{2}$ ; (ii)  $-x, y, -z + \frac{1}{2}$ ; (iii)  $x, -y + 1, z + \frac{1}{2}$ ; (iv)  $x, -y + 2, z - \frac{1}{2}$ .

Data collection: *APEX2* (Bruker, 2007); cell refinement: *APEX2*; data reduction: *SAINT* (Bruker, 2007); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *PLATON* (Spek, 2003); software used to prepare material for publication: *WinGX* publication routines (Farrugia, 1999) and *PLATON*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HK2527).

### References

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## **supplementary materials**

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## 2-Carboxyanilinium chloride monohydrate

**S. A. R. A. Zaidi, M. N. Tahir, J. Iqbal and M. A. Chaudhary**

### Comment

Anthranilic acid is widely used for various purposes such as production of dyes, pigments and saccharin. Its derivatives also have importance in medicinal chemistry and are used as antiinflammatory and anticancer agents (Congiu *et al.*, 2005) and for inhibition of Hepatitis C NS5B polymerase (Nittoli *et al.*, 2005). The title compound has been prepared to see its bioactivity with hope that it will be a good antibacterial agent at the economical cost and to utilize for preparing further derivatives. Crystal structures of proton transfer compound of 2-aminobenzoic acid with nitric acid (Bahadur *et al.*, 2007) has been reported, where the intramolecular N-H···O hydrogen bond has also been observed.

In the molecule of the title compound (Fig. 1), the bond lengths (Allen *et al.*, 1987) and angles are within normal ranges. Ring A (C2-C7) is, of course, planar. The intramolecular N-H···O hydrogen bond (Table 1) results in the formation of a nonplanar six-membered ring B (N1/O2/C1/C2/C7/H1A) having total puckering amplitude,  $Q_T$ , of 0.127 (3) Å, and flattened boat conformation [ $\phi = -21.25$  (3)°,  $\theta = 44.41$  (3)°] (Cremer & Pople, 1975).

In the crystal structure, intermolecular O-H···O and N-H···Cl hydrogen bonds (Table 1) link the molecules (Fig. 2), in which they may be effective in the stabilization of the structure. The C=O···π contact (Table 1) between the carbonyl moiety and the centroid of the benzene ring may further stabilize the structure.

### Experimental

For the preparation of the title compound, 2-aminobenzoic acid (4.11 g, 3 mmol) and trichloroacetic acid (1.64 g, 1 mmol) were neutralized with Na<sub>2</sub>CO<sub>3</sub> separately in H<sub>2</sub>O, and then mixed together. Na<sub>2</sub>CO<sub>3</sub> (3.18 g) was added to the resulting mixture and refluxed for 4 h. The refluxed solution was cooled, and concentrated HCl was added to get PH = 1. The solution was kept in open air for 3 d to get the suitable crystals.

### Refinement

H atoms were located in difference syntheses and refined as [O-H = 0.82 (2) Å (for OH); O-H = 0.76 (3) and 0.78 (3) Å (for H<sub>2</sub>O); N-H = 0.88 (2)-0.958 (19) Å, ( for NH<sub>3</sub>) and C-H = 0.93 (2)-0.96 (2) Å (for CH)] and constrained to ride on their parent atoms with U<sub>iso</sub>(H) = xU<sub>eq</sub>(C,N,O), where x = 1.5 for NH<sub>3</sub> H and x = 1.2 for all other H atoms.

# supplementary materials

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## Figures

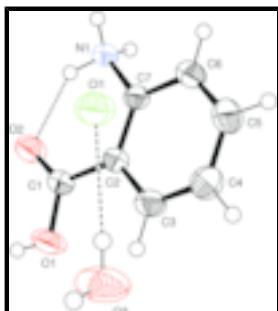


Fig. 1. The molecular structure of the title molecule, with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. Hydrogen bonds are shown as dashed lines.

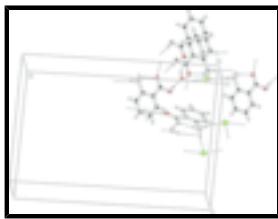


Fig. 2. A packing diagram of the title compound. Hydrogen bonds are shown as dashed lines.

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### Crystal data

$C_7H_8NO_2^+ \cdot Cl^- \cdot H_2O$	$F_{000} = 800$
$M_r = 191.61$	$D_x = 1.407 \text{ Mg m}^{-3}$
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
Hall symbol: -C 2yc	$\lambda = 0.71073 \text{ \AA}$
$a = 23.094 (4) \text{ \AA}$	Cell parameters from 2238 reflections
$b = 4.7833 (8) \text{ \AA}$	$\theta = 2.5\text{--}28.3^\circ$
$c = 16.381 (3) \text{ \AA}$	$\mu = 0.39 \text{ mm}^{-1}$
$\beta = 91.605 (9)^\circ$	$T = 296 (2) \text{ K}$
$V = 1808.8 (5) \text{ \AA}^3$	Prism, brown
$Z = 8$	$0.28 \times 0.10 \times 0.06 \text{ mm}$

### Data collection

Bruker Kappa APEXII CCD diffractometer	2238 independent reflections
Radiation source: fine-focus sealed tube	1749 reflections with $I > 2\sigma(I)$
Monochromator: graphite	$R_{\text{int}} = 0.025$
Detector resolution: 7.5 pixels $\text{mm}^{-1}$	$\theta_{\text{max}} = 28.3^\circ$
$T = 296(2) \text{ K}$	$\theta_{\text{min}} = 2.5^\circ$
$\omega$ scans	$h = -30 \rightarrow 30$
Absorption correction: multi-scan (SADABS; Bruker, 2005)	$k = -6 \rightarrow 3$
$T_{\text{min}} = 0.975, T_{\text{max}} = 0.985$	$l = -21 \rightarrow 20$
9747 measured reflections	

## *Refinement*

Refinement on $F^2$	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.036$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.101$	$w = 1/[\sigma^2(F_o^2) + (0.0476P)^2 + 1.0872P]$ where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.05$	$(\Delta/\sigma)_{\max} < 0.001$
2238 reflections	$\Delta\rho_{\max} = 0.30 \text{ e \AA}^{-3}$
139 parameters	$\Delta\rho_{\min} = -0.22 \text{ e \AA}^{-3}$
Primary atom site location: structure-invariant direct methods	Extinction coefficient: ?

## *Special details*

**Geometry.** Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

**Refinement.** Refinement of  $F^2$  against ALL reflections. The weighted  $R$ -factor  $wR$  and goodness of fit  $S$  are based on  $F^2$ , conventional  $R$ -factors  $R$  are based on  $F$ , with  $F$  set to zero for negative  $F^2$ . The threshold expression of  $F^2 > \sigma(F^2)$  is used only for calculating  $R$ -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement.  $R$ -factors based on  $F^2$  are statistically about twice as large as those based on  $F$ , and  $R$ -factors based on ALL data will be even larger.

## *Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C11	0.05634 (2)	0.81018 (10)	0.08328 (3)	0.0425 (2)
O1	0.24332 (5)	0.7928 (3)	0.36801 (9)	0.0551 (5)
O2	0.17708 (5)	0.9236 (3)	0.45555 (8)	0.0493 (4)
O3	0.18909 (7)	0.6690 (4)	0.07558 (14)	0.0796 (7)
N1	0.07438 (6)	0.6617 (3)	0.45470 (9)	0.0340 (4)
C1	0.19199 (7)	0.7781 (3)	0.39900 (10)	0.0338 (5)
C2	0.15328 (6)	0.5699 (3)	0.35784 (9)	0.0304 (4)
C3	0.17259 (8)	0.4219 (4)	0.29012 (10)	0.0408 (5)
C4	0.13710 (9)	0.2318 (4)	0.25017 (11)	0.0462 (6)
C5	0.08157 (8)	0.1841 (4)	0.27706 (12)	0.0466 (6)
C6	0.06160 (7)	0.3269 (4)	0.34388 (11)	0.0398 (5)
C7	0.09727 (6)	0.5188 (3)	0.38354 (9)	0.0300 (4)
H1	0.2619 (10)	0.922 (5)	0.3885 (14)	0.0661*
H1A	0.0991 (9)	0.802 (4)	0.4684 (12)	0.0510*
H1B	0.0399 (10)	0.732 (4)	0.4423 (13)	0.0510*
H1C	0.0724 (9)	0.534 (4)	0.4996 (12)	0.0510*
H3	0.2113 (9)	0.452 (4)	0.2717 (11)	0.0489*
H3A	0.1584 (14)	0.692 (7)	0.0892 (19)	0.0955*

## supplementary materials

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H3B	0.1945 (14)	0.777 (6)	0.041 (2)	0.0955*
H4	0.1511 (9)	0.133 (5)	0.2061 (13)	0.0554*
H5	0.0582 (9)	0.052 (4)	0.2493 (12)	0.0558*
H6	0.0245 (9)	0.298 (4)	0.3628 (12)	0.0477*

### Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
Cl1	0.0314 (2)	0.0499 (3)	0.0463 (3)	-0.0004 (2)	0.0035 (2)	-0.0055 (2)
O1	0.0320 (7)	0.0668 (9)	0.0673 (9)	-0.0189 (6)	0.0174 (6)	-0.0255 (7)
O2	0.0401 (7)	0.0554 (8)	0.0531 (7)	-0.0144 (6)	0.0133 (6)	-0.0206 (6)
O3	0.0408 (8)	0.0837 (13)	0.1153 (15)	0.0249 (8)	0.0207 (9)	0.0514 (11)
N1	0.0255 (7)	0.0362 (8)	0.0406 (8)	-0.0012 (6)	0.0081 (5)	-0.0030 (6)
C1	0.0283 (8)	0.0362 (8)	0.0372 (8)	-0.0030 (6)	0.0049 (6)	0.0006 (7)
C2	0.0279 (7)	0.0312 (8)	0.0323 (7)	-0.0017 (6)	0.0041 (6)	0.0009 (6)
C3	0.0357 (9)	0.0456 (10)	0.0416 (9)	-0.0040 (8)	0.0111 (7)	-0.0063 (8)
C4	0.0512 (11)	0.0487 (10)	0.0391 (9)	-0.0031 (8)	0.0077 (8)	-0.0135 (8)
C5	0.0445 (10)	0.0462 (10)	0.0486 (10)	-0.0088 (8)	-0.0048 (8)	-0.0109 (8)
C6	0.0292 (8)	0.0435 (10)	0.0466 (9)	-0.0053 (7)	0.0018 (7)	-0.0030 (8)
C7	0.0278 (7)	0.0307 (8)	0.0315 (7)	0.0005 (6)	0.0031 (6)	0.0014 (6)

### Geometric parameters ( $\text{\AA}$ , $^\circ$ )

O1—C1	1.304 (2)	C2—C3	1.399 (2)
O2—C1	1.216 (2)	C2—C7	1.393 (2)
O1—H1	0.82 (2)	C3—C4	1.377 (3)
O3—H3B	0.78 (3)	C4—C5	1.387 (3)
O3—H3A	0.76 (3)	C5—C6	1.380 (3)
N1—C7	1.463 (2)	C6—C7	1.383 (2)
N1—H1B	0.88 (2)	C3—H3	0.96 (2)
N1—H1C	0.958 (19)	C4—H4	0.93 (2)
N1—H1A	0.91 (2)	C5—H5	0.94 (2)
C1—C2	1.487 (2)	C6—H6	0.93 (2)
Cl1···O3	3.1452 (18)	N1···O2	2.6820 (19)
Cl1···N1 <sup>i</sup>	3.3227 (16)	C1···C3 <sup>iv</sup>	3.581 (2)
Cl1···N1 <sup>ii</sup>	3.1464 (16)	C1···O3 <sup>vi</sup>	3.339 (2)
Cl1···N1 <sup>iii</sup>	3.1231 (16)	C1···C4 <sup>iv</sup>	3.477 (3)
Cl1···H5 <sup>iv</sup>	2.96 (2)	C3···C1 <sup>x</sup>	3.581 (2)
Cl1···H3A	2.42 (3)	C3···O1 <sup>viii</sup>	3.338 (2)
Cl1···H6 <sup>v</sup>	3.13 (2)	C4···C1 <sup>x</sup>	3.477 (3)
Cl1···H1B <sup>ii</sup>	2.28 (2)	C1···H1A	2.46 (2)
Cl1···H1A <sup>i</sup>	2.84 (2)	H1···H3B <sup>vi</sup>	2.27 (4)
Cl1···H1C <sup>iii</sup>	2.181 (19)	H1···H3A <sup>vi</sup>	2.27 (4)
O1···C3 <sup>vi</sup>	3.338 (2)	H1···O3 <sup>vi</sup>	1.73 (2)
O1···O3 <sup>vi</sup>	2.539 (2)	H1A···C1	2.46 (2)
O2···O3 <sup>vii</sup>	2.777 (3)	H1A···O2	1.91 (2)

O2···N1	2.6820 (19)	H1A···Cl1 <sup>vii</sup>	2.84 (2)
O3···O2 <sup>i</sup>	2.777 (3)	H1B···Cl1 <sup>ii</sup>	2.28 (2)
O3···O1 <sup>viii</sup>	2.539 (2)	H1B···H6	2.47 (3)
O3···C1 <sup>viii</sup>	3.339 (2)	H1C···Cl1 <sup>ix</sup>	2.181 (19)
O3···Cl1	3.1452 (18)	H3···O1	2.372 (19)
O1···H4 <sup>vi</sup>	2.86 (2)	H3···O1 <sup>viii</sup>	2.655 (19)
O1···H3 <sup>vi</sup>	2.655 (19)	H3A···O2 <sup>i</sup>	2.90 (3)
O1···H3	2.372 (19)	H3A···H1 <sup>viii</sup>	2.27 (4)
O2···H3A <sup>vii</sup>	2.90 (3)	H3A···Cl1	2.42 (3)
O2···H3B <sup>vii</sup>	2.04 (3)	H3B···H1 <sup>viii</sup>	2.27 (4)
O2···H1A	1.91 (2)	H3B···O2 <sup>i</sup>	2.04 (3)
O3···H1 <sup>viii</sup>	1.73 (2)	H4···O1 <sup>viii</sup>	2.86 (2)
N1···Cl1 <sup>ix</sup>	3.1231 (16)	H5···Cl1 <sup>x</sup>	2.96 (2)
N1···Cl1 <sup>vii</sup>	3.3227 (16)	H6···Cl1 <sup>xi</sup>	3.13 (2)
N1···Cl1 <sup>ii</sup>	3.1464 (16)	H6···H1B	2.47 (3)
C1—O1—H1	110.7 (16)	C3—C4—C5	120.07 (17)
H3A—O3—H3B	107 (3)	C4—C5—C6	120.20 (17)
C7—N1—H1C	109.9 (12)	C5—C6—C7	119.54 (15)
H1A—N1—H1B	109.5 (18)	N1—C7—C2	121.21 (13)
H1B—N1—H1C	110.9 (19)	C2—C7—C6	121.35 (14)
C7—N1—H1A	107.7 (13)	N1—C7—C6	117.43 (13)
H1A—N1—H1C	109.0 (17)	C2—C3—H3	119.5 (11)
C7—N1—H1B	109.8 (14)	C4—C3—H3	119.7 (11)
O1—C1—O2	123.05 (15)	C5—C4—H4	120.6 (13)
O1—C1—C2	113.64 (14)	C3—C4—H4	119.3 (13)
O2—C1—C2	123.31 (15)	C4—C5—H5	118.7 (13)
C3—C2—C7	118.04 (14)	C6—C5—H5	121.1 (13)
C1—C2—C7	122.14 (13)	C5—C6—H6	121.5 (12)
C1—C2—C3	119.82 (14)	C7—C6—H6	119.0 (12)
C2—C3—C4	120.80 (17)		
O1—C1—C2—C3	-2.8 (2)	C3—C2—C7—N1	178.91 (14)
O1—C1—C2—C7	178.32 (14)	C3—C2—C7—C6	0.3 (2)
O2—C1—C2—C3	176.32 (16)	C2—C3—C4—C5	-0.3 (3)
O2—C1—C2—C7	-2.5 (2)	C3—C4—C5—C6	0.1 (3)
C1—C2—C3—C4	-178.82 (16)	C4—C5—C6—C7	0.3 (3)
C7—C2—C3—C4	0.1 (2)	C5—C6—C7—N1	-179.16 (16)
C1—C2—C7—N1	-2.2 (2)	C5—C6—C7—C2	-0.5 (3)
C1—C2—C7—C6	179.17 (15)		

Symmetry codes: (i)  $x, -y+2, z-1/2$ ; (ii)  $-x, y, -z+1/2$ ; (iii)  $x, -y+1, z-1/2$ ; (iv)  $x, y+1, z$ ; (v)  $-x, y+1, -z+1/2$ ; (vi)  $-x+1/2, y+1/2, -z+1/2$ ; (vii)  $x, -y+2, z+1/2$ ; (viii)  $-x+1/2, y-1/2, -z+1/2$ ; (ix)  $x, -y+1, z+1/2$ ; (x)  $x, y-1, z$ ; (xi)  $-x, y-1, -z+1/2$ .

#### *Hydrogen-bond geometry ( $\text{\AA}$ , $^\circ$ )*

$D\cdots H$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
O1—H1···O3 <sup>vi</sup>	0.82 (2)	1.73 (2)	2.539 (2)	171 (2)
N1—H1A···O2	0.91 (2)	1.91 (2)	2.6820 (19)	142.1 (18)

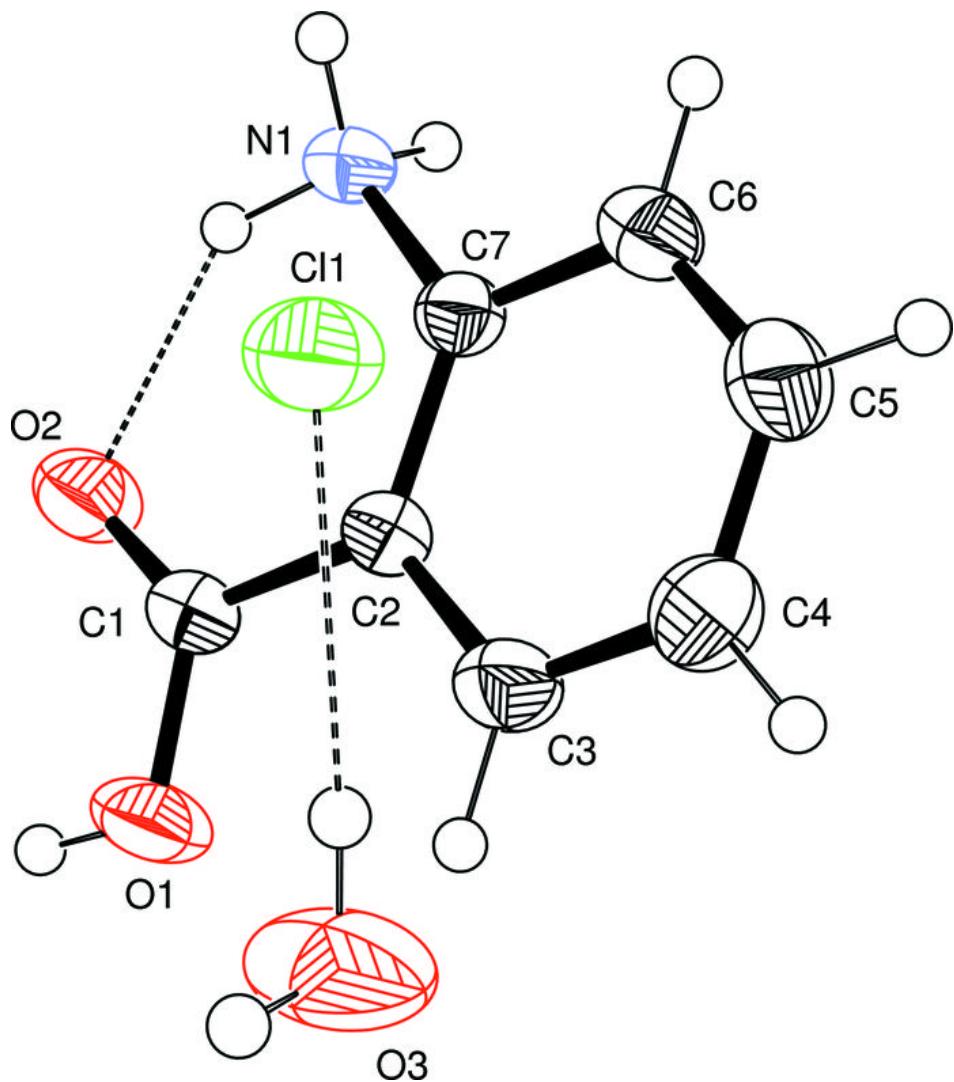
## supplementary materials

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N1—H1B···Cl1 <sup>ii</sup>	0.88 (2)	2.28 (2)	3.1464 (16)	166.6 (18)
N1—H1C···Cl1 <sup>ix</sup>	0.958 (19)	2.181 (19)	3.1231 (16)	167.5 (18)
O3—H3A···Cl1	0.76 (3)	2.42 (3)	3.1452 (18)	160 (3)
O3—H3B···O2 <sup>i</sup>	0.78 (3)	2.04 (3)	2.777 (3)	159 (3)
C1—O2···Cg <sup>iv</sup>	1.216 (2)	3.3988 (16)	3.5802 (18)	88.5 (1)

Symmetry codes: (vi)  $-x+1/2, y+1/2, -z+1/2$ ; (ii)  $-x, y, -z+1/2$ ; (ix)  $x, -y+1, z+1/2$ ; (i)  $x, -y+2, z-1/2$ ; (iv)  $x, y+1, z$ .

Fig. 1



## **supplementary materials**

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**Fig. 2**

